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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

7: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

9: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

9: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

10: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

12: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

13: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

14: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

15: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

16: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

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18: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

19: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

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20: /cgn2_6/ptodata/1/pubpaa/US10B_PUB.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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## ALIGNMENTS

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Publication No. US20030087294A1

GENERAL INFORMATION:
APPLICANT: Wing-Hui WEI

TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC

TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES

TITLE OF INVENTION: THEREOF

FILE REFERENCE: CL001305 DIV

CURRENT APPLICATION NUMBER: US/10/277,032

CURRENT FILING DATE: 2002-10-22

CURRENT FILING DATE: 2002-10-22

CURRENT FILING DATE: 2002-10-22

CURRENT FILING DATE: 2002-10-22
                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FRANCEQ for Windows Version
SEQ ID NO 2
LENGTH: 449
TYPE: PRT
ORGANISM: HomoSapien
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Best Local Similarity
                                                                                                                                                                                                                                            Matches 449; Conservative
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Sequence 2, Application US/10681223

Publication No. US20040081999A1

GENERAL INFORMATION:
APPLICANT: Ming-Hui WEI

TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND
TITLE OF INVENTION: THEREOP

FILE REFERENCE: CL001305 DIV-II
CURRENT APPLICATION NUMBER: US/10/681,223

CURRENT FILING DATE: 2003-10-09

PRIOR APPLICATION NUMBER: 10/277,032

PRIOR APPLICATION NUMBER: 09/984,880

PRIOR APPLICATION NUMBER: 09/984,880

PRIOR APPLICATION NUMBER: 09/984,880

RIOR FILING DATE: 2001-10-31

NUMBER OF SEQ ID NOS: 4

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LENGTH: 449

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                                                                                                     DGRRLQVGCAQVVPVPEPPLHPVVPDLPSSVVFPDREAARAVLEECTSFIPEARAVLDLV
                                                                                                                      DGRRLQVGCAQVVPVPEPPLHPVVPDLPSSVVFPDREAARAVLEECTSFIPEARAVLDLV 240
                                                                                                                                                        GGQAGGAQQGFLLRDPLDDPDTRQALLELLGACQEAPRPHLGEFEADPRGQLWQRLWEVQ
                                                                                                                                                                                                                 DQCPKQIQKGKFQVVAIEGLDATGKTTVTQSVADSLKAVLLKSPPSCIGQWRKIFDDEPT
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                                                                                                                                                                                                                                                                                                                                                                                                     sapiens
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Sequence 4, Application US/10681223
Publication No. US20040081999A1
GENERAL INFORMATION:
APPLICANT: MING-Hui WEI
TITLE OF INVENTION: ISOLATED HUMAN
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Sequence 4, Application US/10277032

Publication No. US20030087294A1

GENERAL INFORMATION:
APPLICANT: Ming-Hui WEI

TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NU
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KIN
TITLE OF INVENTION: THEREOP
FILE REFERENCE: CL001305 DIV
CURRENT APPLICATION NUMBER: US/10/277,032

CURRENT APPLICATION NUMBER: 09/984,880
PRIOR APPLICATION NUMBER: 09/984,880
PRIOR FILING DATE: 2001-10-31
NUMBER OF SEQ ID NOS: 4
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 508
TYPE: PAT
ORGANISM: HomoSapien
US-10-277-032-4
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US-10-277-032-4
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Best Local Similarity 48.1%;
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361
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                                                                    TYAIATEVSGGIQHLPPAHHPVYQWPEDLLKPDLILLLTVSPEERLQRLQGRGMEKTREE 397
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AEAIATEV
                                   AELEANSV 405
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Pred. No. 9.8e-62;
0; Mismatches 4;
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ISOLATED HUMAN KINASE PROTEINS, NUCLEIC

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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(5205)8
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT APPLICATION NUMBER: US/0/360,039
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 911
LENGTH: 196
/ TYPE: PRT
/ ORGANISM: Archaeoglobus fulgidus
US-10-369-493-911
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TITLE OF INVENTION: THEREOF
FILE REFERENCE: CLOO1305 DIV-II
CURRENT APPLICATION NUMBER: US/10/681,223
CURRENT FILING DATE: 2003-10-09
PRIOR APPLICATION NUMBER: 10/277,032
PRIOR APPLICATION NUMBER: 09/984,880
PRIOR FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: 09/984,880
PRIOR FILING DATE: 2001-10-31
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                         Sequence 911, A
Publication No.
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SEQ ID NO 4
LENGTH: 508
TYPE: PRT
ORGANISM: Homo sapiens
S-10-681-223-4
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nes 177; Conservative
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q. US20030233675A1
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Pred. No. 9.8e-62;
0; Mismatches 4
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TITLE OF INVENTION: Identification of Essential Genes in FILE REFERENCE: ELITRA.034A
CURRENT APPLICATION NUMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR PELLING DATE: 2000-03-21
PRIOR PELLING DATE: 2000-03-21
PRIOR PELLING DATE: 2000-05-23
PRIOR PELLING DATE: 2000-05-23
PRIOR PELLING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR PILING DATE: 2000-05-36
PRIOR PELLING DATE: 2000-05-66
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR APPLICATION NUMBER: 60/230,337
PRIOR PILING DATE: 2000-09-06
PRIOR APPLICATION NUMBER: 60/230,347
PRIOR PILING DATE: 2000-10-23
PRIOR PILING DATE: 2000-10-23
PRIOR PILING DATE: 2000-11-27
PRIOR PILING DATE: 2000-11-27
PRIOR PILING DATE: 2000-11-22
PRIOR PILING DATE: 2000-12-22
PRIOR PILING DATE: 2000-12-22
PRIOR PILING DATE: 2000-12-22
PRIOR PILING DATE: 2000-12-76
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US-10-282-122A-67367
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US-10-282-122A-67367
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Best Local Similarity 28.6
Matches 61; Conservative
                                                                                                               SEQ ID NO 67367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 67367, Application US/10282122A Publication No. US20040029129A1 GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
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                                                                                                                                   Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 78614
SOFTWARE: PatentIn version 3.1
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                             ORGANISM: Pasteurella
                                                                                  LENGTH: 243
                                                                                                                                                                                                                        APPLICATION NUMBER: 60/269,308 FILING DATE: 2001-02-16
                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/267,636 FILING DATE: 2001-02-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  308 SLGNYIVASEI-AKES-----AKSPVIVDRYWHSTATYAIATEVSGGLQHLPPAHHPVY 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  163 ETTVVVDASKPLEEVKEEVRKVIESFLNLKKNS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      420 PGCHVVDASPSRE----KVLQTVLSLIQNS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             60 EL--FLKDREIDARENILPALQSGYAVVMDRYYFSNIAYQSARGIDARL---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMNEKIAPKPDLTILLDVEPEIALERVRKRGKLSPFEKLD-----YLRKVRKCFLENAD 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yamamoto, Robert Forsyth, R. Xu, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zamudio, Carlos
Malone, Cheryl
Haselbeck, Robert
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Zyskind, Judith
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Trawick, John
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28.6%; Pred. No. 2.3e-05;
ative 37; Mismatches 75;
                             multocida
                                                                                                                                                                                              See File Wrapper or PALM.
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Query Match

5.98;

Score 140;

DB 15;

Length 243;

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FILE REFERENCE: ELITAL.034
CURRENT APPLICATION NUMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR PRIOR PAPELICATION NUMBER: 60/206,848
PRIOR PRILING DATE: 2000-05-23
PRIOR PRILING DATE: 2000-05-23
PRIOR PRILING DATE: 2000-05-26
PRIOR PRILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR PRILING DATE: 2000-09-06
PRIOR FILING DATE: 2000-09-06
PRIOR PRILING DATE: 2000-09-06
PRIOR PRILING DATE: 2000-09-06
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                                                                                                                                      Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 78614
SOFTWARE: PatentIn version 3.1
SEQ ID NO 53917
LENGTH: 205
        Matches
                          Query Match
Best Local
                                                                                   ORGANISM: Cornyebacterium diptheriae -10-282-122A-53917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 53917; Application US/10282122A Publication No. US20040029129A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/267,636
PRIOR APPLICATION NUMBER: 60/267,636
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PRIOR APPLICATION NUMBER: 60/267,636
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                                                                                                                        TYPE: PRT
                        Local
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                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/269,308
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/230,347
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  412 MSYQRM--ENPGCHVVDASPSREKVLQTVLSLIQ 443
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    51;
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                        Similarity
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Malone, Cheryl
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Zyskind, Judith
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      Conservative
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io, Carlos
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5.6%; Score 131.5; DB 15; 22.4%; Pred. No. 0.003; ative 39; Mismatches 69;
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2; Mismatches 89;
                                                                                                                                                                                                           See File Wrapper or PALM
  Indels
                                      Length 205;
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  69;
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  Gaps
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GENERAL INFORMATION:

APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
ITILE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN ITILE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 11256
LENGTH: 199
TYPE: PRT
                                                             US-10-437-963-196428
                                                                                   RESULT 9
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US-10-369-493-11256
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Sequence 196428, Application US/10437963 Publication No. US20040123343A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 11256, Application US/10369493 Publication No. US20030233675A1
                                                                                                                                              158 ---FLQGVRAIFLKLAADDPERFVVIDASRSPEYIEKEVVKKI
                                                                                                                                                                                                                                   102
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                                                                                                                                                                                                                                                                                                                                                 302 IRRAFYSLGN-----YIVASEIAKESAK-----SPVIVDRYWHSTATYAIATEVSG 347
                                                                                                                                                                                                                                                                                                                                                                                                                         249 KGKFQVVAIEGLDATGKTTVTQSVADSLKAVLLKSPPSCIGQWRKIFDDEPT-----I 301
                                                                                                                                                                                   403 NSVFROKVEMSYORM--ENPGCHVV-DASPSREKVLOTVLSLI 442
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 394 TREEAELEANSVFRQKVEMSYQRMENPGCHY-VDASPS--REKVLQTV 438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       58;
                                                                                                                                                                                                                                                                                                                                                                                               1 RGK--LITLEGIDGSGKSTVAEK------LQKNPE--IKAFKPVFTREPTRGTLTGDA 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            55 TDSAYAMATLFALDRYDAKAVLGRYVGTSKV-----VLLDRYVASNAAYSAAR----
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                                                                                                                                                                                                                               GITLKTRLENPL-EWVKDLHRSWTIVPDLTFLFDIRPEISIERCGKRGEQSKFEKLE---
                                                                                                                                                                                                                                                                    GLQHLPPAHHPVYQWPEDL-----LKPDLILLTVSPEERLQRLQGRGMEKTREEAELEA 402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -----TRNDAMVQWVQELEFEELGLPVPDIHILLHTSPELAAQRAQRREATDASRKLDR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 128.5; DB 15; Pred. No. 0.0053; 9; Mismatches 71;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 15; Length 199;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  55;
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei

Tue

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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Coldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10(52052)B
CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
FRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 21577
LENGTH: 205
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                                                                                                                                                 ; ORGANISM: Pyrococcus abyssi
US-10-369-493-21577
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; OTHER INFORMATION: Clone ID: PAT_MRT4530_92280C.1.pep
US-10-437-963-196428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 196428
LENGTH: 253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21577, Application US/10369493 Publication No. US20030233675A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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Best Local Similarity
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                                                                        y Match 5.4%; Score 127.5; DB 15; Local Similarity 22.7%; Pred. No. 0.0068; hes 52; Conservative 43; Mismatches 71;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             378
                                          249
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                           KGKFQVVAIEGLDATGKTTVTQSVADSLKAVLLKSPPSCIGQWRKIFDDEPT-----II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LKELAMSCILE 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SPEERLQRLQGRGMEKTREEAELEANSVFRQKVEMSYQRMENPGCHVVDASPSREKVLQT 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -AKSPVIVDRYWHSTATYAIATEVSGGLQHLPPAHHPVYQW---PED-LLKPDLILLLTV 377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EGWR--FPDRGT-----SVGQ-MISAYLANESELDDRTIHLLFSANRWEKRALMERKL 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LPOWRGVFRSVRMESGSSQGGRGALIVLEGLDRSGKSSQCARLLSFLQG-----KGCQA 82
     RGYF--VVLEGIDGSGKTTQAKLLAEWFEE-----QGWDVLLTKEPTDTEFGRLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QPEKAAER-GGYGGER-YEKIE-----FQKKVGEHYHSLRDSTWKVVDGSLPMEVVEEQ 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LGGTTLIVDRYSYS----GVAFSAAKGLD-----IEWCKAPENGLIAPDLVVYLDV 179
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US-10-425-114-64336
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US-10-369-493-22800
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Sequence 64336, Application US/10425114
Publication No. US20040034888A1
GENERAL INFORMATION:
APPLICANT: Liu, Jingdong
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SEQ ID NO 22800
LENGTH: 259
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Schizosaccharomyces pombe
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                                                                                                                                               221
                                                                                                                                                                                                                      169
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ilarity 22.3%;
Conservative 4
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APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Marry S.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN TITLE OF INVENTION: MINBER: US/10/369,493
CURRENT APPLICATION NUMBER: US/10/369,493
PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR FILING DATE: 2003-02-21
PRIOR FILING DATE: 2002-02-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(259)
; OTHER INFORMATION: unsure at all xaa locations
US-10-369-493-22800
                                                                                                                                                                                                363 - PEDILKPDLILLITVSPEERIQRLQGRGMEKTREEAELEANSVFRQKVEMSYQRME---
                                                                                                                                                                                                                                                                     118 AFSGIAFSAAKVILSPLGXFVNYRRLRIXTLXTIYEVTNIKXGLD-------WEWCKS 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          248 QKGKFQVVAIEGLDATGKTTVTQSVADSL-----KAVLLKSP------
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  285 --PSCIGQWRKIFD------DEPTIIRRAFYSLGNYIVASEIAKESAKS-PVIVDRY
                                                                                                                                                                                                                                                                                                                                                                                                63 DRTTAIG--KKIDDYLKESVQLNDQ---VIHLLFSANRWETIQYIYEQINKGVTCILDRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 KONRGRLIVÍEGÍÐRSGKSTOCÓLLVÐKLISOHEKAELFKFÞGKSYFCFLADXLVTFTRX 62
REEGLEFITLDASSSLEDVHSQIVDLVSN 249
                                                             -NPGCHVV--DASPSREKVLQTVLSLIQN 444
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                                                                                                                                     PDRGLPRPDLVIFLNVDP--RIAATRGQYGEERYEKIEMQ---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 127; DB 1
Pred. No. 0.01;
42; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 15;
                                                                                                                                                                                                                                                                                                                                  ----TEVSGGLQHLPPAHHPVYQW----
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                                                                                                                                                                                                                                                                                                                                                                                                    117
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With FILLE OF INVENTION: Plants and Uses Thereof for Plant Improvement CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
LENGTH: 263
TYPE: PDT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-815-242-11027
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ORGANISM: Zea mays

FEATURE:

OTHER INFORMATION: Clone ID: LIB3632-055-C3_FLI.pep
US-10-425-114-64336
                                                                       PRIOR FILING DATE: 2000-05-23

PRIOR APPLICATION NUMBER: 60/20
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/24
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/25
PRIOR APPLICATION NUMBER: 60/25
PRIOR FILING DATE: 2000-11-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11027, Application US/09815242 Patent No. US20020061569A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
                                                                                                                                                                                                                                                    FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Habelbeck, Robert APPLICANT: Ohlsen, Kari L. APPLICANT: Zyskind, Judith W.
                                                                                                                                                                                                                                                                                                                 APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification
TITLE OF INVENTION: Prokaryotes
                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                    APPLICATION NUMBER: 60/242,578
FILING DATE: 2000-10-23
APPLICATION NUMBER: 60/253,625
FILING DATE: 2000-11-27
APPLICATION NUMBER: 60/269,308 FILING DATE: 2001-02-16
                                                APPLICATION NUMBER: 60/257,931
                                                                                                                                                               APPLICATION NUMBER: 60/207,727
                                      FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                     Yamamoto, Robert T.
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                                    2000-12-22
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Pred. No. 0.012;
4; Mismatches 70; Indels 71;
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                                                                                                                                                                                                                                                                                                                                Essential Genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        70;
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PRIOR FILING DATE: 2000-09-09
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/267,636
PRIOR APPLICATION NUMBER: 60/267,636
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US-10-282-122A-58176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11027
LENGTH: 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 58176, Application US/10282122A Publication No. US20040029129A1
Remaining Prior Application
NUMBER OF SEQ ID NOS: 78614
SOFTWARE: PatentIn version 3
SEQ ID NO 58176
                                                                        PRIOR APPLICATION NUMBER: 60/2
PRIOR FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/2
PRIOR FILING DATE: 2001-02-16
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Identification of Essential Genes in Microorganisms FILE REFERENCE: ELITRA.034A CURRENT APPLICATION NUMBER: US/10/282,122A CURRENT FILING DATE: 2003-02-20 PRIOR APPLICATION NUMBER: 60/191,078 PRIOR PILING DATE: 2000-03-21 PRIOR APPLICATION NUMBER: 60/206,848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
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PRIOR FILING DATE: 2000-09-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     : Wang, ... Zamudio, Carryl Pob
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                        PatentIn version 3.1
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Wall, Daniel
Trawick, John
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Yamamoto, Robert
Forsyth, R.
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Haselbeck, Robert
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Pred. No. 0.0095;
Pred. No. 0.0095;
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PRIOR APPLICATION NUMBER: US 60/360,039
PRIOR PILING DATE: 2002-02-21
NUMBER OF SED ID NOS: 47374
SEQ ID NO 1320
LENGTH: 205
TYPE: PRT
ORGANISM: Pyrococcus horikoshii
US-10-369-493-1320
Search completed: February 27, 2005, 17:59:13 Job time : 134 secs
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US-10-369-493-1320
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1320, Application US/10369493 Publication Nd. US20030233675A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 5.3%; Score 126; DB 15; Length 210; Best Local Similarity 26.1%; Pred. No. 0.0095; Matches 57; Conservative 32; Mismatches 91; Indels 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Glater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: doldman, Barry S.
APPLICANT: Chen, Xianfeng
TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052) B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/369,493
CURRENT FILING DATE: 2003-02-28
                                                                                                                                   400 LEANSVFRQKVEMSYQRMEN--PGCHVVDASPSREKVLQTVLSLIQN 444
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                                                                                         FDKIVELQRKVRHNYLKLAEMFPEMRIVNALSSIEDIHSDIVALVKH 200
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

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Ratent No. 6489153

GENERAL INFORMATION:

APPLICANT: Ming-Hui WEI

TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC

TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES

TITLE OF INVENTION: THEREOF

FILE REFERENCE: CL001305

CURRENT APPLICATION NUMBER: US/09/984,880

CURRENT FILING DATE: 2001-10-31

NUMBER OF SEQ ID NOS: 4

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 2

LENGTH: 449

TYPE: PRT

ORGANISM: HomoSapien

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17308, A	24702, A	7769, Ap	25512, A	32145, A	30625, A	20357, A	31851, A	<ol> <li>Appli</li> </ol>	30034, A	8122, Ap	12293, A	<ol><li>Appli</li></ol>	4, Appli	6, Appli	8, Appli	25366, 7	46, Appl

## ALIGNMENTS

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QWPEDLLKPDLILLLTVSPEERLQRLQGRGMEKTREEAELEANSVFRQKVEMSYQRMENP
                                     DQCPKQIQKGKFQVVAIEGLDATGKTTVTQSVADSLKAVLLKSPPSCIGQWRKIFDDEPT
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; Sequence 4, Application U
; Patent No. 6489153
; GENERAL INFORMATION:
; APPLICANT: Wing-Hui WEI
; TITLE OF INVENTION: ISOL
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SOFTWARE: FastSEQ for Windows Version 4.0
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Patent No. 6664087
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; TITLE OF INVENTION: THEREOF
; FILE REFERENCE: CL001305
; CURRENT APPLICATION NUMBER: US/09/984,880
; CURRENT FILING DATE: 2001-10-31
; NUMBER OF SEQ ID NOS: 4
; SOFTMARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 4
; SEQ ID NO 4
; ENGTH: 508
                                                                                          ; ORGANISM: HomoSapien US-10-277-032-4
                                                                                                                                                                          APPLICANT: Ming-Hai WEI
TITLE OF INVENTION: ISOLATED HUMAN KINASE PRITITLE OF INVENTION: ACID MCLECULES ENCODING
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL001305 DIV
CURRENT APPLICATION NUMBER: US/10/277,032
CURRENT FILING DATE: 2002-10-22
PRIOR APPLICATION NUMBER: 09/984,880
PRIOR FILING DATE: 2001-10-31
NUMBER OF SEQ ID NOS: 4
NUMBER OF SEQ ID NOS: 4
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US-10-277-032-4
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; LENGTH: 508
; TYPE: PRT
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                                 Matches
                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                Sequence 4, Application Patent No. 6664087
GENERAL INFORMATION:
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                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version
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ECTSFIPEARAVLDLVDQCPKQIQKGKFQVVAIEGLDATGKTTVTQSVADSLKAVLLKSP
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                              0,
                            Score 795.5; DB 4
Pred. No. 2.9e-72;
0; Mismatches 4
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Pred. No. 2.9e-72;
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                                                          DB 4;
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                              Indels
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CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR PILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 28002
LENGTH: 956.
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US-09-252-991A-28002
US-09-252-991A-28002, Application US/09252991A
; Sequence 28002, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION: Rubenfield et al.
; APPLICANT: MARC J. Rubenfield et al.
APPLICANT: MUCLEIC ACID AND A
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Best Local Similarity
Matches 124; · Conserv
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
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  MNDADSLQDYQRVRQQAIRSLFEIVEGSSEGTVIVDREARIVWINQRYAQRFGLADPAQA
                                          VPDLPS--
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                                                                                                                                                                                                                                                                                                                  AGAVPPRLRAADPAALGDLLRQPDPALLHGQLAALAAPGKRPDTQCGEPGDLDVPLRRHP
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                                                                               RRARRA---TDPPGPLARRPGAA--GKRRRAGRRGILSLHRERLPGYSGGPFRRPPQEPP
                                                                                                                                                                                                FORCQLLRLLCYCPGGQAG---GAQQGFLLRDPLDDPDTRQALL-----ELLGACQEA 156
                                                                                                                                                                                                                                                                         ---DPRLA----ALLGPP-----ERSYSLCVPVTPDAGCGARVRAARLHQRLLHQLRRGP
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                                                                                                                                                           PER---LRQQLLPGPGTRHRHRLGAGRGPLRFDP--RPTVRQPAAGHAHIRGEHLHACGDS
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TITLE OF INVENTION: Crystallization and Structure of Invention: Staphylococcus Aureus The FILE REFERENCE: 6245.NCP
CURRENT APPLICATION NUMBER: US/09/632,553
CURRENT FILING DATE: 1090-08-04
PRIOR APPLICATION NUMBER: 60/147,117
PRIOR FILING DATE: 1399-08-04
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; SEQ ID NO 3
; LENGTH: 216
; TYPE: PRT
; ORGANISM: Saccharomyces cerevisiae
US-09-632-553-3
                                                                                                                                    RESULT 7
US-09-252-991A-25690
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US-09-632-553-3
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Best Local S
Sequence 25690, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AEROGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                  164
                                                                                                                                                                                                                                                                                                111 MDLDWCLQPDVGLLKPDLTLFLSTQDVDNNAEKSGFGDERYE--
                                                                                                                                                                                                                                                                                                                                  358 PVYQW---PE-DILKPDLILLTVSPEERLQRLQGRGMEKTREEAELEANSVFRQKVEMS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           222 VLEECTSFIPEA-----RAVLDLVDQCPKQIQKGKFQVVAIEGLDATGKT--TVT
                                                                                                                                                                                                                                                                                                                                                                           61 SDQAIHLLFSANRWEIVDKIKKDLLEGKNIVMDRYVYSGVAYSAAKGTNG---
                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 GRGKLILIEGLDRTGKTTQCNILYKKLQPNCKLLKFPERSTRIGGLINEY--LTDDSFQL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                FMXLLDKEIRKGDESITIVDVT---NKGIQEVEALI
                                                                                                                                                                                                                                                       YQ-----RMENPGCHVVDASPSREKVLQTVLSLI 442
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RRGAA----SEAPVILLIGETGTGKELLAHAIHSASPRAHKAFVSVNAAAIPESLLEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmacia & Upjohn
Pharmacia & Upjohn
VENTION: Crystallization and Structure Determination
VENTION: Cranhvlococcus Aureus Thymidylate Kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5.0%;
ilarity 25.0%;
Conservative 3:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 118.5; DB 4;
Pred. No. 0.0011;
17; Mismatches 80;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         216;
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US-09-632-553-2 .
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CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 25690
LENGTH: 681
TYPE: PRT
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                  SOFTWARE: PA
SEQ ID NO 2
LENGTH: 213
TYPE: PRT
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                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Pharmacia & Upjohn
TITLE OF INVENTION: Crystallization and Structure Dete
TITLE OF INVENTION: Staphylococcus Aureus Thymidylate
FILE REFERENCE: 6245.NCP
CURRENT APPLICATION NUMBER: US/09/632,553
CURRENT FILING DATE: 2000-08-04
                                                                                                                                                                                                                                                                                        Sequence 2, Application US/09632553
Patent No. 6689595
                                                                                                               CURRENT FILING DATE: 2000-08-04
PRIOR APPLICATION NUMBER: 60/147,117
PRIOR FILING DATE: 1999-08-04
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Best Local Similarity
Matches 125; Conserv
                                                                                               NUMBER OF SEQ ID NOS:
ORGANISM: Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                             510
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             221
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51 APGDADAPDPRLAALLGPPERSYSLCVPVTPDAGC--GARVRAARLHQRLLHQLRRGPFQ 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97
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                                                                           PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                   LDEALRAEPLVOWORKOASILACOOOLORLEOWCDRCLGELAPSC----OPLREVIAO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KSPVIVDRYWHSTATYAIATEVSGGLQ------HLPPAHHPVYQ-----W---PE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EPÍDPGQPCGPSLRYDPDYDRLRELRREDDSSLPTGVWQÁEAKRADWAAVEQLÁSDLLQR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 D-----Likedlill-----LTVSPEERLQRLQGRGMEKTREEAELEANSVFRQK 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RSKDLMLAAWLGEAW--LQRGGLGGLQRALVLLAELCERYPEEVHPQAQDGDQSWRVPPI 449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAI-----AVRDAV--AGHRROHGRYPMT----LPLSGN-----ALSLEVLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AVLEECTSFIPEARAVLDLVDQCPKQIQKGKFQVVAIEGLDATGKTTVTQSVADSLKAVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RLHGVSAGPATAGVCALOPGKR----GADVPP-----PVVDEPGRQDPAVHPAAAGAR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGARRAGPGSLAGP-AGRRGLRPPPGAGAALARPAVRARQVGPARRRPGHRLAQ----RP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QPGTRRRPGDRLEPVLGRPGKPFPRRAGQGPDLECRRSAENHAALGEERLAASGRRP---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5.0%; Score 118; DB 4; L
23.2%; Pred. No. 0.0072;
vative 49; Mismatches 189;
                                                                                                                                                                                                                                                                                                                                                                                                                                        EMSYORMEN------PGCHVVDASPSREKVLQ 436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 4; Length 681;
                                                                                                                                                                                                                             Determination
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RESULT 10
US-09-902-540-15299
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US-09-489-039A-7540
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                                                                             Sequence 15299, Application Patent No. 6833447 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: PUBLICACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: PUBLICACION DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT FILING DATE: 2709.489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 7540
LENGTH: 219
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 52; Conserv
APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 7540, Application US/09489039A Patent No. 6610836
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Klebsiella pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                   165 IEQESMNFFNRTRARYLELA---AADPSIRTVDATQPLDAVARDIRATIAQWMAE
                                                                                                                                                                                                                                                                                                                                         340
                                                                                                                                                                                                                                                                                                                                                                                                           289 GQWRKIFDDEPTIIRRAFYSLGNYIVASEJAKESAKSP-VIVDRYWHSTATY-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         248 QKGKFQVVAIEGLDATGKTTVTQSVADSLKA-----VLLKSPPSCI------
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 116 D--QHMLATLRDAVLGDFRPDLTLYLDVTPEVGLKKARARGELDRIEQESFDFFNRTRAR 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            409 KVEMSYQRMENPGCHVVDASPSREKVLQTVLSLIQNSFSE 448
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                                                                                                                                                                                                                                                                                                                                                                                                                                                5 KKMRSNYIVIEGLEGAGKTTARQLVVETLQSAGIHDMVFTREPLGTILAEKLRSLVLDIQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             64 VITDKAEVL--MFYAARVQLVETVİKPALANGTWVİGDRHDLSTQAY-----QGGGRGI 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 IVÍÉGLEGAGKTTARNVVVETLEQLGIRDMVFTREÞGGTQLAEKLRSLVLDIKSVGD--E
                                                                                                                                                                                                                                                   TREEAELEANSVFRQKVEMSYQRMENPGCHVVDASPSREKVLQTVLSLIQNSFSE 448
                                                                                                                                                                                                                                                                                                RTMLATLRDAVLGD-----
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                                                                                                                                                                                                                                                                                                                                 ----AIATEVSGGLQHLPPAHHPVYQWPEDLIKPDLILLTVSPEERLQRLQGRG-MEK 393
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5.0%; Score 117.5; DB 22.1%; Pred. No. 0.0014;
                                                                                                                    US/09902540
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                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              85;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 219;
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                                                                                                                                                                                                                   216
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RESULT 11
US-09-902-540-15011
; Sequence 15011, Application US/09902540
hatent No. 6833447
  APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Serill Reference: 38-10(15849)B
FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Myxococcus xanthus US-09-902-540-15299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 15299
LENGTH: 925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
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37; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGEVVPVSLCERLLARVP-ARLHHQYGPTE-----ATIDASCWDCVPGETRSYVPMG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DGGKASAVPEDEWPSSAASEPYPLTPMQEGMLFHTLLAPGSRVYHEQLEFELRGPLV---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TPTGKVDRGALPEPSEF---NRLVPSGLHEAPRGALEERLSRLWGEVLRVALPGRHAHFL 380
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VVVRGEGEDEHRL---VAYVVPARNAVEPLELTALRAFLERSLPDYMVPAVFIPLDGLPL 323
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Pred. No. 0.024;
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US-09-107-433-3820
US-09-107-433-3820
; Sequence 3820, Application US/09107433
; Patent No. 6800744
; GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
; APPLICANT: Lynn A Doucette ACID AMINO ACID
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
THERAPEUTICS
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US-09-902-540-15011
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SEQ ID NO 15011
LENGTH: 1646
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COMPUTER: «Unknown»

OPERATING SYSTEM: «Unknown»

SOFTWARE: «Unknown»

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/107,433

FILING DATE: 30-Jun-1998
                                                                                                                         ZIP: 02354
COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM IS09660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  --APRPHLGEFEADPRGQLWQ------RLWEVQDGRRLQVGCAQVVPVPEPPLHPV 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ----GAQQGFLLR-----DPLDD-PDTR-----QALLELLGACQE---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CV-PVTPD----AGCGARVRAARLHQRLLHQLRRGPFQRCQLLRLLCYCPGGQAG---
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                                                                                                                                                                                           STATE: Massachusetts COUNTRY: USA
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                                                                                                                                                                                                                                       CITY: Waltham
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Pred. No. 0.17;
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Sequence 2424, Application US/09252991A

Sequence 2424, Application US/09252991A

Patent No. 6551795

GENERAL INFORMATION:

APPLICANT: Marc J: Rubenfield et al.

TITLE OF INVENTION: AUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 24254

LENGTH: 955

TYPE: PRI

TYPE: PRI

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TYPE: PRI

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                 Matches
                                                                     Query Match
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Best Local
                                                                                                                                           ORGANISM: Pseudomonas aeruginosa
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TELEFAX: (781)893-8277
INFORMATION FOR SEQ ID NO: 3820:
                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
les 53; Conserv
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LOCATION: (B) LOCATION 1...218
SEQUENCE DESCRIPTION: SEQ ID NO: 3820:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein HYPOTHETICAL: YES ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-011
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/ 085131
FILING DATE: May 12, 1998
APPLICATION NUMBER: 60/051553
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           171 LHKKVRQGYLSLLDKEGNRIVKIDASLPLEQVVETTKAVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               124
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           74;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           64 SHTQMDAKTELLLYIASRRQHLVEKVLPALEAGKLVIMDRFIDSSVAYQGFGRGLDIBAI 123
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TYPE: amino acid
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           Conservative
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  4.6%; Score 109; DB 4; Length 955; 28.1%; Pred. No. 0.1; ative 14; Mismatches 83; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.6%; Score 109; DB. 24.1%; Pred. No. 0.01; tive 36; Mismatches
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     92;
  Gaps
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US-09-252-991A-25642
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US-09-252-991A-25642
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PRIOR FILING DATE: 1998-07-27
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 25642
LENGTH: 615
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: MARG J. Rubenfield et al.
APPLICANT: MARG J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196,136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 25642, Application US/09252991A Patent No. 6551795
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ORGANISM: Pseudomonas aeruginosa
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       350
                                                                                                      293
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                                                                                                                                                                                                                                                                                        177 WEVQDGRRLQVGCAQVVPVPEPPLHPVVPDLPSSVVFPDREAARAVLEECTSFIPEARAV 236
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                                                                                                                                                                                              237 LDLVDQCPKQIQKGKFQVVAIEGLDATGKTTVTQSVADSLKAVLLKSPPSCI----GQWR
                                                                                                                                                                                                                                                                                                                                                                                            122 GOAGGAQQGFLLRDPLDDPDTRQALLELLGACQE--APR---PHLGEFEADPRGQLWQRL 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    280
                                                                                                                                                                                                                                                                                                                                                                                                                                             326 ELADLELPEALPLLIAALRGDPDPGVRGEAARLLE-----AWEEDAVVDALCAA--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           66 ----LGPPERSYSLCVPVTPDAGCGARVRAARLHQRLLHQLRRGPFQRCQLLRLLCYCPG 121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6 RILRGPLSGPLLGRRGVCAGAMAPPCRFVLELPDCTLAHFALGADAPGDADAPDPRLAAL 65
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
QHLPPAHHPVYQWPEDLLKPDLILLLTVSPEERLQRLQGRGMEKTREEAELEANSVFRQK 409
                                                                                               KIFDDEPTIIRRAFYSLGNYIVASEIAKESAKSPV---IVDRYWHSTATYAIATEVSGGL
                                                -----VREEAATTLG-----KLGREEAGEPLLKALADDYWQVRLR---AARALGRL
                                                                                                                                                                                                                                                      REL----RLEESAA----
                                                                                                                                                LGWLRHQPALAELAR---LASADVDPEVRRAATGALGLSREATVL--PALCAALADAQWQ 506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RLAAGHHPGPTADRHMTCKNATLPE---ILEL-----SPRLEDADPGVRRLALI 325
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Pred. No. 0.057;
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                                                                                                                                                                                                                                                 ----PALAALGDPQAA---
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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FILE REFERENCE: GM10201
CURRENT APPLICATION NUMBER: US/09/259,109
CURRENT FILING DATE: 1999-02-26
NUMBER OF SEG ID NOS: 3
SOFTWARE: FadtSEQ for Windows Version 3.0
SEQ ID NO 2
LENGTH: 212;
TYPE: PRT
ORGANISM: Streptococcus pneumoniae
US-09-259-109-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/09259109
Patent No. 6270762
GENERAL INFORMATION:
APPLICANT: Burnham, Martin K. R.
APPLICANT: Zalacain, Magdalena
APPLICANT: Chalker, Alison F.
APPLICANT: Chalker, Alison F.
APPLICANT: Ingraham, Karen A.
APPLICANT: Traini, Christopher M.
APPLICANT: Warren, Patrick V.
TITLE OF INVENTION: tdk
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Search completed: February 27, 2005, 17:53:15 Job time : 46 secs
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                                                                                                                414 Y-QRMENPGCHVV--DASPSREKVLQTVLSLI 442
                                                                                                                                                                                                                                                                         300 TIIRRAFYSLGNYIVASEIAKESAKSPYIVDRYWHSTATY----AIATEVSGGLQHLPP 354
                                                                                                                                                                                                                                                                                                                585 SAQALRVAEG-----DGDPEVRKAVRIALAQLR 612
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550 RHRP----
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                                                                                                                                                                                         355 AHHPVYQWPEDLLKPDLILLITVSPEERLQRLQGRGMEKTREEAELEANSV-FRQKVEMS 413
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                                                                               173
                                                                                                                                                                                                                                    66 TELLLYIASRROHLVEKVLPALEAGKLVİMDRFIDSSVAYQGFGRGLDIEAIDWLN---- 121
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YLSTLDKEGNRIVKIDASLPLEQVVETTKAVL 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90; Indels 36;
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
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10
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2276
2125.2
1694.2
1579.6
1579.6
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1: /cgn2_6/ptodata/2/pubpna/VS07 PUBCOMB.seq:*
2: /cgn2_6/ptodata/2/pubpna/PCT NEW PUB.seq:*
3: /cgn2_6/ptodata/2/pubpna/VS06 NEW PUB.seq:*
4: /cgn2_6/ptodata/2/pubpna/VS06_PUBCOMB.seq:*
5: /cgn2_6/ptodata/2/pubpna/VS08_PUBCOMB.seq:*
6: /cgn2_6/ptodata/2/pubpna/VS08_PUBCOMB.seq:*
6: /cgn2_6/ptodata/2/pubpna/VS08_PUBCOMB.seq:*
7: /cgn2_6/ptodata/2/pubpna/VS09B_PUBCOMB.seq:*
9: /cgn2_6/ptodata/2/pubpna/VS09B_PUBCOMB.seq:*
10: /cgn2_6/ptodata/2/pubpna/VS09B_PUBCOMB.seq:*
11: /cgn2_6/ptodata/2/pubpna/US09C_PUBCOMB.seq:*
12: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*
13: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*
14: /cgn2_6/ptodata/2/pubpna/US10B_PUBCOMB.seq:*
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2571
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10713.331 Million cell updates/sec
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               US-10-277-032-1
US-10-681-223-1
US-10-305-810-6
US-10-305-810-720
US-10-370-715B-645
US-10-357-930-20954
US-10-357-930-22065
US-10-357-930-22065
US-10-357-930-220800
US-10-357-930-27924
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               Sequence 1, Appli
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Sequence 720, App
Sequence 645, App
Sequence 20954, A
Sequence 22065, A
Sequence 26800, A
Sequence 27924, A
Sequence 3, Appli
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2.8	2.8			2.8		_	2.9	2.9	2.9	2.9	2.9	3.0	3.0	3.0	3.0	3.0	3.0	3.1	3.1	4.9	5.8	8 5	9.3	11.8	14.2		15.2	15.5	16.0	16.0	16.0	16.0	40.3
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US-10-616-187-48	US-10-023-523-48	US-10-023-529-48	US-09-976-740-48	US-10-437-963-8597	US-10-156-761-5988	US-10-156-761-1	US-10-156-761-1558	US-10-425-115-164510	US-10-767-701-9739	US-10-723-860-5759	US-10-767-701-4633	US-10-267-255-76	US-09-953-348-76	US-10-267-255-51	US-09-953-348-51	US-10-729-802-1	US-10-425-115-17653		US-10-156-761-7362	US-09-783-590-11496	US-10-240-425-827	US-09-783-590-5564	US-10-357-930-1516	US-09-867-701-2642	US-10-723-860-5376	US-10-357-930-31877	US-10-357-930-3179	US-10-357-930-10685	US-10-357-930-12348	US-10-357-930-42442	US-10-357-930-40840	-10-357-	US-10-264-237-104
48,	48,	Sequence 48, Appl	Sequence 48, Appl	8597,	Sequence 5988, Ap		Sequence 1558, Ap	164510	9739,	5759,	4633	76,	76,	51,	13	Sequence 1, Appli	Sequence 17653, A	Sequence 1, Appli	Sequence 7362, Ap	Sequence 11496, A	Sequence 827, App	Sequence 5564, Ap	Sequence 1516, Ap	2642, A	Sequence 5376, Ap		3179,		12348,	Sequence 42442, A	Sequence 40840, A	33519	Sequence 104, App

## ALIGNMENTS

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PRIOR APPLICATION NUMBER: 09/984,880
PRIOR FILING DATE: 2001-10-31
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.(
SEQ ID NO 1
LENGTH: 2571
TYPE: DNA
ORGANISM: HomoSapien
US-10-277-032-1
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Publication No. US20030087294A1

GENERAL INFORMATION:
APPLICANT: Ming-Hui WEI
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL001305 DIV
                                                                                                                     Query Match 100.
Best Local Similarity 100.
Matches 2571; Conservative
                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/277,032
CURRENT FILING DATE: 2002-10-22
                 61 GCCACTGTCGGGGCGCTCGCTCGGGCGCGGGGTCTGGGGGCCATGGCTCCGCC
61
                                                          100.0%; Score 2571;
100.0%; Pred. No. 0;
vative 0; Mismatches
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2281 AATGCAATGTTTGTGTATTTTCATTACACAAACTTAATTTGTCTTGTTAAATAAGTT 2340	Ş	1201 CATGGAGAAGACCAGGGAAGAAGCAGAACTTGAGGCCAACAGTGTGTTTCGTCAAAAGGT 1
2221 AATATGTTTCCTCAGCATAGGAGCTATGATTAATTAAAAAGTGGAGTCAAAACGCTA 2280	D QQ	CCTTATCCTGCTGCTCACTGTGAGTCCTGAGGAGAGGGTTGCAGAGGCTGCAGGGCCGGGG
2161 TCTGAACTGGATGCATATCATTTAGCCAATCCAGTAATTTATTT	B &	GCACCTGCCCCAGCCCATCACCCTGTGTACCAGTGGCCAGAGGACCTGCTCAAACCTGA 1
2101 ATTTATATTTCTTTTTTTCTAAAGAATTGGCTCTGAATAGAATGCACATTTTCCA 2160	B &	<u> </u>
41	g Qy	961 CAATTATATTGTGGCCTCCGAAATAGCTAAAGAATCTGCCAAATCTCCTGTGATTGTAGA 1020
	dg Qy	
1921 TGCCCAAGGGACACTCCTGCAGAGCCATTTTTGGGTAAGGAACACTTACAAAGAAGGCAT 1980 	B 8	841 GTCAGTGGCAGATTCACTTAAGGCTGTCCTCTTAAAGTCACCACCCTCTTGCATTGGCCA 900
1861 CCCTCAGTCCCAAATTCTAGTAGCCAAGTGTTCCTGCAGAGGCTGTCTATGTGTCCTGGC 1920	B &	Y 781 AAAGTTCCAGGTTGTTGCCATCGAAGGACTGGATGCCACGGGTAAAACCACGGTGACCCA 840
ATGCCACATTGGCTTCTTCCACATCTGGTTGCACTCGTCATGATGGGCTCGCTGCATCT	g Q	y 721 TCCTGAAGCCCGGCAGTGCTTGACCTGGTCGACCAGTGCCCAAAACAGATCCAGAAAGG 780
CACCAGGAAGACCCCGGAGCCTCTGCTTACACTGGCATGCTGATGAGATCGTGAC 	D Qy	y 661 CGTGGTCTTCCCGGACCGGGAAGCCGCCCGGGCCGTTTTGGAGGAGTGTACCTCCTTTAT 720
AGACCTGTCTTACTCAGACAGCACCAGGTGTACAGAGCAGCAGAAGATCTTCCAGAT	D Q	y 601 ACAGGTCGTGCCCGGAGGCCCCGCTGCACCCGGTGGTGCCAGAGTTGCCCAGTTC 660
IGGCATAGTAAACATTTTCTTGGAACATATGTTTCACTTAATCACTACCAAATATCTGGA 	. B &	y 541 CCAGCTGTGGCAGCGCCTCTGGGAGGTGCAAGACGGCAGGCGGCTGCAGGTGGGCTGCGG 600
GTCATCACAAAGGTCCCTCTCATCATGTTCCAGTGAGAGGCCAGGATTGCTTTCTTCC	dg dg	y 481 GGGCGCCTGCCAGGAGGCACCACGCCCGCACTTGGGCGAGTTCGAGGCCGACCCGCGCGG 540
CTGGAGACCAGGCATCTTAATTTTACTTCAGCCATCGTACCCTCTTCTGACTGA	da	y 421 CTTCCTGCTGCGGGACCCCCTGGATGACCCCTGACACCCGGCAAGCCTGCTCGAGCTGCT 480
41 ACCATTIGITATGCAGTGTTCCCAAATTTCTGTTCTACAGCAGGAGTGTGGTGGCAGAAAA	g dy	y 361 CCAGCTGCTCAGGCTGCTCTGCTACTGCCCGGGCGGCCAGGCCGGCGGCACAGCAAGG 420
GTTACTCTGGCCAGGTGCCACGTCTAACTAGATTAGATGTTGTATGAAACATCTACATCC	ρ	y 301 GGCGGCGCGCTGCACCAGCGCCTGCTGCACCAGCTGCGCCGCGGCCCCTTCCAGCGGTG 360
21 CAGAGAAAAGGTCCTGCAGACAGTATTAAGCCTAATCCAGAATAGTTTTAGTGAACCGTA 	QY db	y 241 GCGCAGCTACTCGCTGTGCCGTGACCCCGGACGCCGGCTGCGGGGCCCGGGTCCG 300
61 AGAAATGTCCTACCAGCGGATGGAGAATCCTGGCTGCCATGTGGTTGATGCCAGCCCTC	₽ \$	181 CGCCCCGGCGACGCAGACGCCCCGACCCCCGCCTGGCGGCGCTGTTGGGGGCCCCCGGA 240
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Qy 361 CCAGCTGCTCAGGCTGCTCTGCTACTGC	Qy 301 GGCGGCGCGGCTGCACCACGCCTGCTG	OY 241 GCGCAGCTACTCGCTGTGCGTGCCCGTG	Qy         181 CGCCCCGGGGACGCAGACGCCCCCGAC	Qy 121 GTGCCGCTTCGTCCTGGAGCTTCCCGAC	Qy 61 GCCACTGTCGGGGGCCGCTGCTCGGGCGG	Qy 1 cggggccgAgggcggcgTcgCTgAggcgcccATggc 	Query Match 100.0%; Score 2571; Best Local Similarity 100.0%; Pred. No. 0; Matches 2571; Conservative 0; Mismatches		PRIOR FILING DATE: 2001-10-31  NUMBER OF SEQ ID NOS: 4  SOFTWARE: FASTSEQ for Windows Version 4.	CURRENT FILING DATE: 2003-10-09 PRIOR APPLICATION NUMBER: 10/277,032 PRIOR FILING DATE: 2002-10-22 PRIOR APPLICATION NUMBER: 09/984 880	; TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, ; TITLE OF INVENTION: THEREOF ; PILE REPERENCE: CLOO1305 ENCOLOS 305	Publication Nd. US20040081999A1 GENERAL INFORMATION: APPLICANT: Ming-Hui WEI TITLE OF INVENTION: ISOLATED HUMAN KIND	) RESULT 2 US-10-681-223-1 ; Sequence 1. Application US/10681223	Qy 2521 AAGCCTTGCATGTCCTGCAGCGTCGCACTAAAGGAGGGGGCTTTTGCACCC	Qy 2461 ATTATAGAGCTTGTGTGACTTTAGGGAC	Qy 2401 TACATATGAAATCGCTTTGACATATCTT	Qy 2341 CAAGTGGATCTTGGAGTGGATTTCTTC	2281
CCAGCTGCTCAGGCTGCTCTGCTACTGCCCGGGCGGCCAGGCCAGGCGGCGCGCGC	GGCGGCGCGCTGCACCAGCGCTGCTGCACCAGCTGCGCCGCGGCCCCTTCCAGCGGTG 360	GCGCAGCTACTCGCTGTGCGTGCCCGTGACCCCGGACGCCGGCTGCGGGGGCCCGGGTCCG 300	CGCCCCGGCGACACACACCCCCAACCCCCGCCTGGCGGCGCTGTTGGGGCCCCCGGA 240	GTGCCGCTTCGTCCTGGAGCTTCCCGACTGCACCCTGGCTCACTTCGCCCTAGGCGCCGA 180	GCCACTGTCGGGGCCGCTGCTCGGGCGGCGCGGGGTCTGCGCTGGGGCCATGGCTCCGCC 120	CTTCGCCCGCCGGCTCCTGCGCGG 60	571; DB 17; Length 2571; o. 0; tches 0; Indels 0; Gaps 0;		.0		ODING HUMAN KINASE PROTEINS, AND USES	SE PROTEINS. NICLEIC		TAAAGGAGGGGCTTTTGCACCC 2571 	ATTATAGAGCTTGTGTGACTTTAGGGAGTAGCTGTCTTATACACATACTCAAGCCCTG 2520 	TACATATGAAATCGCTTTGACATATCTTTAGACAGAAAAAGTAGCTGAGTGAG	CAAGTGGATCTTGGAGTGGGATTTCTTGGTAAATTATCTTGCACTTGAATGTCTCATGAT 2400 	AATGCAATGTTTGTTGTGTATTTTCATTACACAAACTTAATTTGTCTTGTTAAATAAGTT 2340
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1441 ACCATTTGTTATGCAGTGTTCCCAAATTTCTGTTCTACAAGCATGTTGTGTGGGGAGAAAA 1500 	1381 GTTACTCTGGCCAGGTGCCAGGTCTAACTAGATTAGATGTTGTTTGAAACATCTACATCC 1440 1381 GTTACTCTGGCCAGGTGCCACGTCTAACTAGATTAGATGTTTGTT	21 CASACAMAMOSTIC ISCAGACASIANI ANGCETAMI CASACATANG ITITANG SEMACESIA I (ASCAGACASIANI TANGCETAMI CASACATANG ITITANG SEMACESIA I (ASCAGACAGACAGACAGACAGACAGACAGACAGACAGACA	bl AGAMATETCCTACCAGCGGATGGAGATCCTGGCTGCCATGTGGTTGATGCCAGCCCCTC 61 AGAMATGTCCTACCAGCGGATGGAGAATCCTGGCTGCCATGTGGTTGATGCCAGCCCCTC 62 AGAMATGTCCTACCAGCGGATGGAGAATCCTGGCTGCCATGTGGTTGATGCCAGCCCCTC	01 CATGGAGAAGACCAGGGAAGAAGCAGAACTTGAGGCCAACAGTGTGTTTCGTCAAAAGGT [	41 CCTTATTCCTGCTGCTCACTGTGAGGTCCTGAGGAGAGGTTGCAGAGGCCGGGG 	81 GCACCTGCCCCAGCCCATCACCCTGTGTACCAGTGGCCAGAGGACCTGCTCAAACCTGA	1021 CAGGTACTGGCACAGCACGGCCACCTATGCCATAGCCACTGAGGTGAGTGGGGGTCTCCA 1080		901 GTGGAGGAAGATCTTTGATGAACCAACTATCATTAGAAGAGCTTTTTACTCTTTTGGG 960 	841 GTCAGTGGCAGATTCACTTAAGGCTGTCCTCTTAAAGTCACCACCCTCTTGCATTGGCCA 900	781 AAAGTTCCAGGTTGTTGCCATCGAAGGACTGGATGCCACGGGTAAAACCACGGTGACCCA 840	721 TCCTGAAGCCCGGGCAGTGCTTGACCTGGTCGACCAGTGCCCAAAACAGATCCAGAAAGG 780	661 CGTGGTCTTCCCGGACCGGGAAGCCGCCCGGGCCGTTTTGGAGGAGTGTACCTCCTTTAT 720	601 ACAGGTCGTGCCCGGAGCCCCGCTGCACCCGGTGGTGCCAGACTTGCCCAGTTC 660	541 CCAGCTGTGGCAGCGCCTCTGGGAGGTGCAAGACGGCAGGCGGCTGCAGGTGGGCTGCGC 600	481 GGGCGCTGCCAGGAGGCACCACGCCCGCACTTGGGCGAGTTCGAGGCCGACCCGCGCGCG	421 CTTCCTGCTGCGCGACCCCCTGGATGACCCTGACACCCGGCAAGCGCTGCTCGAGCTGCT 480	

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                                              ATTATAGAGCTTGTGTGACTTTAGGGAGTAGCTGTCTCTTATACACATACTCAAGCCCTG
                                                              ATTATAGAGCTTGTGTGACTTTAGGGAGTAGCTGTCTCTTATACACATACTCAAGCCCTG
                                                                                               CAAGTGGATCTTGGAGTGGGATTTCTTGGTAAATTATCTTGCACTTGAATGTCTCATGAT
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                                                                                                                                                CAAGTGGATCTTGGAGTGGGATTTCTTGGTAAATTATCTTGCACTTGAATGTCTCATGAT
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Sequence 6, Application US/10305810

Publication No. US20030176385A1

GENERAL INFORMATION:

APPLICANT: UJU, JIngfang
APPLICANT: Huang, Chunli
APPLICANT: Huang, Chunli
APPLICANT: Tailon, Bruce B.
APPLICANT: Tailon, Bruce B.
APPLICANT: Simons, Jan Fredrik
APPLICANT: Simons, Jan Fredrik
APPLICANT: Tailon, Bruce B.
APPLICANT: Millet, Isabelle
TITLE OF INVENTION. ANTISENSE MODULATION OF PROTEIN EXP
FILE REFERENCE: 21402-501
CURRENT APPLICATION NUMBER: US/10/305,810
CURRENT APPLICATION NUMBER: 60/334,148
PRIOR APPLICATION NUMBER: 60/334,148
PRIOR APPLICATION NUMBER: 60/334,148
PRIOR APPLICATION NUMBER: 60/336,572
PRIOR APPLICATION NUMBER: 60/336,572
PRIOR APPLICATION NUMBER: 60/192,838
PRIOR APPLICATION NUMBER: 60/192,838
PRIOR APPLICATION NUMBER: 60/194,256
PRIOR APPLICATION NUMBER: 60/194,256
PRIOR APPLICATION NUMBER: 60/194,256
PRIOR APPLICATION NUMBER: 60/194,256
PRIOR APPLICATION NUMBER: 60/233,798
PRIOR APPLICATION NUMBER: 60/233,798
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PRIOR APPLICATION NUMBER: 60/233,798
PRIOR APPLICATION NUMBER: 60/233,798
PRIOR APPLICATION NUMBER: 60/233,798
PRIOR APPLICATION NUMBER: 60/240,316
PRIOR FILING DATE: 2000-02-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 60/240,316
PRIOR FILING DATE: 2000-03-15
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; ORGANISM: Thymidine Kinase
US-10-305-810-6
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US-10-305-810-6
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Best Local Similarity
Matches 2388; Conserv
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SEQ ID NO 6
LENGTH: 2856
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TACTGCCCGGGCGGCCAGGCCGGCGCGCACAGGCTTCCTGCTGCGCGACCCCCTG
                                                                              CTGCTGCACCAGCTGCGCCGCCGCCCTTCCAGCCGTGCCAGCTGCTCAGGCTGCTCTGC
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nilarity 98.9%;
Conservative
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RESULT 4 US-10-044-090-720	1519	1460 TCCCAAATTTCTGTTCTACAAGCATGTTGTGTGGGCAGAAAACTGGAGACCAGGCATCTTA
2480 TTTAGGGAGTAGCT 2493 	OY 	1400 ACGTCTAACTAGATTAGATGTTGTTGAAACATCTACATCCACCACTTTGTTATGCAGTGT
	1399	1340 ACAGTATTAAGCCTAATCCAGAATAGTTTTAGTGAACCGTAGTTACTCTGGCCAGGTGCC
GATTICTTGGTAAATTATCTTGCACTTGAATGTCTATGATGATATGACATATGAAATGGCTTTG	2AG 1339 Db	1280 ATGGAGAATCCTGGCTGCCATGTGGTTGATGCCAGCCCCTCCAGAGAAAAAGGTCCTGCAG
ATTTTCATTACACAAACTTAATTTGTCTTGTTAAATAAGTTCAAGTGGATCTTGGAGTGG 2	1279 1194	
2240 GGAGCTATGATTCATTAATTAAAAGTGGAGTCAAAACGCTAAATGCAATGTTTCTTGTGT 2299	9AA 1219 QY     Db	1160 GTGAGTCCTGAGGAGAGGGTTGCAGAGGCTGCAGGGCCGGGGCATGGAGAAGACCAGGGAA 
ATTTAGCCAATCCAGTAATTTATTTATATTAATCTATACATAATATGTTTCCTCAGCATA	1159	1100 CACCCTGTGTACCAGTGGCCAGAGGACCTGCTCAAACCTGACCTTATCCTGCTGCTCACT
2120 TITICTAAAGAATTGGCTCTGAATAGAATGCACATTTTCCATCTGAACTGGATGCATATC 2179	1099	1040 GCCACCTATGCCATAGCCACTGAGGTGAGTGGGGGTCTCCAGCACCTGCCCCAGCCCAT
ACTGCAAATAFACCAACCTTCTCTGAATTATATTTTGCTTATTTATATTTCTTTTCTTT	1039	980 GAAATAGCTAAAGAATCTGCCAAATCTCCTGTGATTGTAGACAGGTACTGGCACAGCACGG
CAGAGCCTTTTGATAGGCTTCTGATGTCATTCATAAAAACATTCAAGCCAAGATGCTCC	979	920 GATGAACCAACTATCATTAGAAGAGCTTTTTACTCTTTGGGCAATTATATTGTGGCCTCC
CAGAGCCATTTTTGGGTAAGGAACACTTACAAAGAAGGCATTGATCTTGTGTCTGAGGCT 199	919	860 AAGGCTGTCCTCTTAAAGTCACCACCCTCTTGCATTGGCCAGTGGAAGAAGATCTTTGAT
GTAGCCAAGTGTTCCTGCAGAGGCTGTCTATGTGTCCTGGCTGCCCAAGGGACACTCCTG	859 780	803 GAAGGACTGGATGCCACGGGTAAAACCACGGTGACCCAGTCAGTGGCAGATTCACTT
CACATCTGGTTGCACTCGTCATGATGGGCTGCATGCATCCTCAGTCCCAAATTCTA	ATC 802 Db	743 GACCTGGTCGACCAGTGCCCAAAACAGATCCAGAAAAGGAAAGTTCCAGGTTGTTGCCATC
CTCTGCTTCTCCTACACTGGCATGCTGATGAGATCGTGACATGCCCACATTGGCTTCTT 1	742	683 GCGCCCGGGCCGTTTTGGAGGAGTGTACCTCCTTTATTCCTGAAGCCCGGGCAGTGCTT
AGCACCAGGTGTACAGAAGCAGCAGACAAGATCTTCCAGATCAGCAGGGAGACCCCGGAG	682	623 CCCCGCTGCACCCGGTGGTGCCAGACTTGCCCAGTTCCGTGGTCTTCCCGGAACCGGGAA
TTGGAACATATGTTTCACTTAATCACTACCAAATATCTGGAAGACCTGTCTTACTCAGAC 	540	563 GRIGGTGCAAGACGGCAGGCGGCTGCAGGTGGGCTGCGCACAGGTCGTGCCCGTGCCCGAG
CTCATCATGITCCAGTGMAGAGCCAGCGATTGCTTCTTCCTGGCATAGTAAAAATITTC	562 480	503 CCCCCGCACTTCGGCGAGTTCGAGGCCGACCCGCCGCCGCCAGCTGTGGCAGCCCTCTGG
ATTITUDE CONTROL CONTR	502 420	443 GATGACCCTGACACCCGGCAAGCGCTGCTCGAGCTGCTGGGGCGCCTGCCAGGAGGCACCA
1375 TCCCCAAATTTCTGTTCTACAAGCATGTTGTGGGCAGAAAACTGGAGACCAGGCATCTTA 1434	360	301 TACTGCCGGGCGAGGCCGGCGCGCACAGCAAGGCTTCCTGCTGCGCGACCCCCTG

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Sequence 720, Application US/10044090
PUDIICATION NO. US20020137081A1
GENERAL INFORMATION:
APPLICANT: Olga Bandman
TITLE OF INVENTION: GENES DIFFERENTIALLY EXPRESSED IN
FILE REFERENCE: PA-0028 US
CURRENT APPLICATION NUMBER: US/10/044,090
CURRENT FILING DATE: 2002-01-09
NUMBER OF SEQ ID NOS: 850
SOFTWARE: PERL PROGRAM
SEQ ID NO 720
LENGTH: 2645
TYPE: DNA
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Best Local :
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FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. US20020137081A1 197362.
5-10-044-090-720
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      GAATCTGCCAAATCTCCTGTGATTGTAGACAGGTACTGGCACAGCACGGCCACCTATGCC
                                                                                      ATCATTAGAAGAGCTTTTTACTCTTTGGGCAATTATATTGTGGCCTCCGAAATAGCTAAA
                                                                                                                                                                 TTAAAGTCACCACCTCTTGCATTGGCCAGTGGAGGAAGATCTTTGATGATGAACCAACT
                                                                                                                                                                                                                    GATGCCACGGGTAAAACCACGGTGACCCAGTCAGTGGCAGATTCACTTAAGGCTGTCCTC
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Query Match Best Local Similarity 98.9%; Pred. No. 0; Matches 1779; Conservative 0; Mismatches 13; Indels 7; Gaps 7;  Qy 695 GTTTTGGAGGAGTGTACCTCCTTTATTCCTGAAGCCCGGGCAGTGCTTGACCTGGTCGAC 754	APPLICANT: SCHOENFELD, JILL R. APPLICANT: SCHOENFELD, JILL R. APPLICANT: WILLIAMS, P. MICKEY APPLICANT: WOOD, WILLIAM I.  APPLICANT: WU, THOMAS D. TITLE OF INVENTION: Compositions and Methods for the Treatment of Immune TITLE OF INVENTION: Related Diseases FILE REFERENCE: P1948R1-US CURRENT APPLICATION NUMBER: US/10/370,715B CURRENT APPLICATION NUMBER: US/10/370,715B CURRENT FILING DATE: 2003-02-21 NUMBER OF SEQ ID NOS: 742 SEQ ID NO 645 LENGTH: 2566 TYPE: DNA ORGANISM: Homo sapien US-10-370-715B-645	ហ ភា 🖰	Qy 2372 AATTATCTTGCACTTGAATGTCTCATGATTACATATGAAATCGCTTTGACATATCTTTAG 2431	Qy 2132 TTGGCTCTGAATAGAATGCACATTTTCCATCTGAACTGGATGCATATCATTTAGCCAATC 2191
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APPLICANT: Endege, Wilson

APPLICANT: Endege, Wilson

APPLICANT: MORDADAN, JOHN

ITITLE OF INVENTION: HOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR

ITITLE OF INVENTION: HOVEL GENES, COMPOSITIONS, KITS, AND THER

ITITLE OF INVENTION: HOWEL GENES, COMPOSITIONS, KITS, AND THER

ITITLE OF INVENTION: HOWEL GENES, COMPOSITIONS, KITS, AND THER

ITITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THER

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US-10-357-930-20954
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RESULT 7

US-10-357-930-22065

US-10-357-930-22065

Sequence 22065, Application US/10357930

Publication No. US20040259086A1

GENERAL INFORMATION:
APPLICANT: Endegel. Robert
APPLICANT: Endegel, Wilson
APPLICANT: Monahan, John
ITITLE OF INVENTION: HOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLE OF INVENTION: HUMAN PROSTATE CANCER

FILE REFERENCE: MRI-007BCN
CURRENT FILING DATE: 2003-02-04
PRIOR APPLICATION NUMBER: US/10/357,930
CURRENT FILING DATE: 2003-02-16
PRIOR APPLICATION NUMBER: 09/785,276
PRIOR PILING DATE: 2003-02-16
PRIOR PILING DATE: 2003-02-16
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR PILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/189,862
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SEQ ID NO 22065
LENCTH: 2199
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: 1, 2, 3, 4, 2198, 2199
OTHER INFORMATION: n = A,T,C or G
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PRIOR APPLICATION NUMBER: 60/207,454
PRIOR FILING DATE: 2000-05-25
PRIOR PELLING DATE: 2000-06-09
PRIOR PILING DATE: 2000-06-09
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR FILING DATE: 2000-07-18
PRIOR APPLICATION NUMBER: 60/255,281
PRIOR FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 62332
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Best Local Similarity 97.1%;
Matches 1748; Conservative
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Sequence 2600, Application US/10357930

| Publication No. US20040259086A1
| GENERAL INFORMATION: Schlegel. Robert APPLICANT: Schlegel. Robert APPLICANT: Schlegel. Robert ITILE OF INVENTION: MOTAL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLE OF INVENTION: HUMAN PROSTATE CANCER FILE REFERENCE: MRI-007BCR
| CURRENT APPLICATION NUMBER: US/10/357,930
| CURRENT APPLICATION NUMBER: US/10/357,930
| CURRENT PILLING DATE: 2003-02-04 |
| PRIOR APPLICATION NUMBER: 09/785,276 |
| PRIOR APPLICATION NUMBER: 60/183,319 |
| PRIOR APPLICATION NUMBER: 60/183,319 |
| PRIOR APPLICATION NUMBER: 60/189,862 |
| PRIOR APPLICATION NUMBER: 60/207,454 |
| PRIOR APPLICATION NUMBER: 60/207,454 |
| PRIOR APPLICATION NUMBER: 60/211,314 |
| PRIOR APPLICATION NUMBER: 60/211,314 |
| PRIOR APPLICATION NUMBER: 60/211,314 |
| PRIOR APPLICATION NUMBER: 60/213,007 |
| PRIOR APPLICATION NUMBER: 60/213,007 |
| PRIOR APPLICATION NUMBER: 60/255,281 |
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SOFTWARE: PASTSEQ for Windows Version
SEQ ID NO 26800
LENGTH: 2199
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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TAATTTATTATATATCATACATAATATGTTTCCTCAGCATAGGAGCTATGATTCAT
                                             AGGCTTCTGATGTCATTAAAGACATTCAAGCCAAGATGCTCCAACTGCAAATATATACC
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Sequence 27934, Application US/10357930

Publication No. US20040259086A1

GENERAL INFORMATION:
APPLICANT: Schlegel. Robert
APPLICANT: Endege, Wilson
APPLICANT: Honosham, John
ITILE OF INVENTION: IDENTIFICATION, KITS, AND METHODS FOR TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND THER TITLE OF INVENTION: HOPEL GENES, COMPOSITIONS, KITS, AND THER TITLE OF INVENTION: HOPEL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLE OF INVENTION: HOPEL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR TITLING DATE: 2003-02-04

PRIOR APPLICATION NUMBER: 60/183,319

PRIOR APPLICATION NUMBER: 60/183,319

PRIOR APPLICATION NUMBER: 60/207,454

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PRIOR APPLICATION NUMBER: 60/211,314

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; LOCATION: 1, 2, 3, 4, 2198, 2
; OTHER INFORMATION: n = A,T,C
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Sequence 3, Application US/10277032

| Sequence 3, Application US/2030087294A1
| GENERAL INFORMATION: US20030087294A1
| GENERAL INFORMATION: HOLATED HUMAN KINASE PROTEINS, NUCLEIC TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND TITLE OF INVENTION: THEREOF
| FILLE REFERENCE: CL001305 DIV | CURRENT APPLICATION NUMBER: US/10/277,032 | CURRENT FILING DATE: 2002-10-22 | PRIOR APPLICATION NUMBER: 09/984,880 | PRIOR PRILING DATE: 2001-10-31 | NUMBER OF SEQ ID NOS: 4 | SOFTWARE: FastSEQ for Windows Version 4.0 | SEQ ID NO 3 | LENGTH: 20966 | Type: DNA | CRGANISM: HomoSapien | Type: DNA | CRGANISM: HomoSapien | TYPE: DNA | CRGANISM: HomoSapien | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA | TYPE: DNA |
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98.1%;
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GTAGCTGAGTGAGGGGGAAATTATAGAGCTTGTGTGTGACTTTAGGGAGTAGCT
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US-10-681-23-3
; Sequence 3, Application US/10681223
; Publication No. US20040081999A1
; GENERAL INFORMATION:
APPLICANT: Ming-Hui WEI
; TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES
; TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL001305 DIV-II
   CURRENT FILING DATE: 2003-10-09
   PRIOR APPLICATION NUMBER: 10/277,032
   PRIOR APPLICATION NUMBER: 09/984,880
   PRIOR APPLICATION NUMBER: 09/984,880
   PRIOR APPLICATION NUMBER: 09/984,880
   PRIOR FILING DATE: 2001-10-31
; SOPTWARE: FastSEQ for Windows Version 4.0
; SOPTWARE: PastSEQ for Windows Version 4.0
; SOPTWARE: Assessed for Windows Version 4.0
; CRGANISM: Homo sapiens
US-10-681-223-3
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Best Local Similarity 98.1%;
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TGATGGGCTCGCTCCTCCCTCAGTCCCAAATTCTAGTAGCCAAGTGTTCCTGCAGAG
                                                       ATGCTGATGAGATCGTGACATGCCCACATTGGCTTCCTCCACATCTGGTTGCACTCGTCA
                                                                                                                             CAGACAAGATCTTCCAGATCAGCAGGGAGACCCCGGAGCCTCTGCTTCTCCTACACTGGC
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                                                                                                                                                                                                 TCACTACCAAATATCTGGAAGACCTGTCTTACTCAGACAGCACCAGGTGTACAGAAGCAG
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APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and
FILE REFERENCE: PA131P1
CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR PTLING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
JUMMER OF SEQ ID NOS: 2876
SOFTWARE: Patentin Ver. 3.1
SEQ ID NO 104
LENGTH: 1579
TYPE: DNA
ORGANISM: Homo sapiens
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US-10-264-237-104
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                                                              Query Match 40.3%;
Best Local Similarity 98.8%;
Matches 1127; Conservative
                                                                                                                                                                                                                                                                                                             Sequence 104, Application US/10264237
Publication No. US20040009491A1
GENERAL INFORMATION:
                          1353 TAATCCAGAATAGTTTTAGTGAACCGTAGTTACTCTGGCCAGGTGCCACGTCTAACTAGA 1412
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                                                               Score 1035.4; DB
Pred. No. 2e-245;
0; Mismatches
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                                                                                     DB 17;
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APPLICANT: Endege, Wilson
APPLICANT: Endege, Wilson
APPLICANT: Monahan, John
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR
ITITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THEF
TITLE OF INVENTION: HUMAN PROSTATE CANCER
FILE REFERENCE: MRI-007BCN
CURRENT APPLICATION NUMBER: US/10/357,930
CURRENT FILING DATE: 2003-02-04
PRIOR FILING DATE: 2003-02-16
PRIOR PRILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR FILING DATE: 2000-02-17
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR APPLICATION NUMBER: 60/213,314
PRIOR APPLICATION NUMBER: 60/211,314
PRIOR APPLICATION NUMBER: 60/211,314
PRIOR APPLICATION NUMBER: 60/211,314
PRIOR PRILING DATE: 2000-06-09
PRIOR PILING DATE: 2000-06-09
PRIOR PILING DATE: 2000-07-18
PRIOR PILING DATE: 2000-07-18
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Best Local Similarity 98.6%;
Matches 415; Conservative
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LENGTH: 454
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SOFTWARE: FastSEQ for Windows Version
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APPLICANT: Endege, Wilson
APPLICANT: MONAHAN, John
ITITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS FOR
ITITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THEF
ITITLE OF INVENTION: HUMAN PROSTATE CANCER
FILLE OF INVENTION: HUMAN PROSTATE CANCER
FILLE REFERENCE: MRI-007BCN
CURRENT FILLING DATE: 2003-02-04
PRIOR APPLICATION NUMBER: 09/785,276
PRIOR FILLING DATE: 2003-02-16
PRIOR FILLING DATE: 2003-02-16
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR APPLICATION NUMBER: 60/183,319
PRIOR APPLICATION NUMBER: 60/213,319
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR FILLING DATE: 2000-03-16
PRIOR APPLICATION NUMBER: 60/207,454
PRIOR FILLING DATE: 2000-06-09
PRIOR APPLICATION NUMBER: 60/211,314
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR APPLICATION NUMBER: 60/219,007
PRIOR APPLICATION NUMBER: 60/219,007
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Best Local Similarity 98.6%;
Matches 415; Conservative
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SOFTWARE: FastSEQ for Windows Version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGTCCTGAGGAGAGGTTGCAGAGGCTGCAGGGCCCGGGCATGGAGAAGACCAGGGAAAGAA 1222
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Sequence 42442, Application US/10357930

| Publication No. US20040259086A1
| GENERAL INFORMATION: APPLICANT: Schlegel. Robert APPLICANT: Schlegel. Robert APPLICANT: Endege, Wilson APPLICANT: Monahan, John ITILE OF INVENTION: INDENTIFICATION, ASSESSMENT, PREVENTION, AND THER TITLE OF INVENTION: HUMAN PROSTATE CANCER FILE REFERENCE: MAI-1007BCN
| CURRENT APPLICATION NUMBER: US/10/357,930 |
| CURRENT APPLICATION NUMBER: 09/785,276 |
| PRIOR APPLICATION NUMBER: 60/183,319 |
| PRIOR APPLICATION NUMBER: 60/183,319 |
| PRIOR APPLICATION NUMBER: 60/183,319 |
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| PRIOR APPLICATION NUMBER: 60/207,454 |
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| PRIOR APPLICATION NUMBER: 60/211,314 |
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| PRIOR APPLICATION NUMBER: 60/219,007 |
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| PRIOR APPLICATION NUMBER: 60/255,281 |
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; ORGANISM: Homo sapiens
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US-10-357-930-42442
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Best Local Similarity 98.6%;
Matches 415; Conservative
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                                           CAAATTTCTGTTCTACAAGCATGTTGTGTGGCAGAAAACTGGAGACCAGGCATCTTAATT
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Search completed: February 28, 2005, 22:36:20 Job time : 1440 secs

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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*

2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*

3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*

4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*

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5 US-09-984-880-3
4 US-09-266-965-51
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5 US-09-103-840A-1
10S-09-616-289-48
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10S-08-642-255-50
10S-08-642-255-50
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10S-09-92-688-8
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10S-09-93-688-8
10S-09-93-688-2
10S-09-430-141
10S-09-647-144A-14
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sequence 1, Appii		Sequence 11, Appl	Sequence 985, App	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 9209, Ap	Sequence 9, Appli	Sequence 193, App	Sequence 1, Appli	Sequence 48, Appl	-	•	Sequence 15, Appl	•	Sequence 15, Appl

ALIGNMENTS

## Sequence 1, Application US/09984880 Fatent No. 6489153 GENERAL INFORMATION: APPLICANT: Ming-Rui WEI TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES TITLE OF INVENTION: THEREOF FILE REFERENCE: CL001305 CURRENT FILLING DATE: 2001-10-31 NUMBER OF SEQ ID NO: 4 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO: 1 LENGTH: 2571 TYPE: DNA ORGANISM: HomoSapien Query Match 100.0%; Score 2571; Best Local Similarity 100.0%; Pred. No. 0; Matches 2571; Conservative 0; Mismatches DB 4. **;** Length Indels 2571; 0 Gaps

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1501 CTGGGGGGGGGTTTTTTTTTTTCTTCAGCCATCGTTACCTTTCTGACTGA

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Sequence 1, Application US/10277032

Sequence 1, Application US/10277032

Patent No. 6664087

GENERAL INFORMATION:

APPLICANT: Ming-Hui WEI

TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC

TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND

TITLE OF INVENTION: THEREOF

FILE REFERENCE: CL001305 DIV

CURRENT FILING DATE: 2002-10-22

PRIOR APPLICATION NUMBER: 09/984,880

PRIOR FILING DATE: 2001-10-31

NUMBER OF SEQ ID NOS: 4

SOFTWARE: FRatSEQ for Windows Version 4.0

SEQ ID NO 1

LENGTH: 2571

TYPE: DNA

ORGANISM: HomoSapien

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RESULT 3
US-09-984-880-3
Sequence 3, Application US/09984880
Patent No. 6489153
GENERAL INFORMATION:
APPLICANT: Ming-Hai WEI
TITLE OF INVENTION: 1SOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CLOO1305
CURRENT APPLICATION NUMBER: US/09/984,880
CURRENT FILING DATE: 2001-10-31
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
ENGTH: 20966
TYPE: DNA
ORGANISM: HomoSapien
US-09-984-880-3
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Query Match 44.0%; Score 1130.4; DB 4; Best Local Similarity 98.1%; Pred. No. 6.3e-262; Matches 1228; Conservative 0; Mismatches 16;
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Sequence 3, Application US/10277032

Patent No. 6644087

GENERAL INFORMATION:
APPLICANT: Ming-Hui WEI
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES
TITLE OF INVENTION: THERBOF
FILE REFERENCE: CL001305 DIV
CURRENT APPLICATION NUMBER: US/10/277,032
CURRENT FILING DATE: 2002-10-22
PRIOR APPLICATION NUMBER: 09/984,880
PRIOR FILING DATE: 2001-10-31
NUMBER OF SEQ.ID NOS: 4
SOFTWARE: FRAUSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 20966
TYPE: DNA
ORGANISM: HomoSapien
US-10-277-032-3
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TCACTACCAAATATCTGGAAGACCTGTCTTACTCAGACAGCACCAGGTGTACAGAAGCAG
                                                 CCAGCGATTGCTTCCTTGCCATAGTAAACATTTTCTTGGAACATATGTTTCACTTAA
                                                                                  ATGTTGTGGCAGAAAACTGGAGACCAGGCATCTTAATTTTACTTCAGCCATCGTACCC
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ilarity 98.1%;
Conservative
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18144 TCACTACCAAATATCTGGAAGACCTGTCTTACTCAGACAGCACCAGGTGTACAGAAGCAG 1820:
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                  GTAGCTGAGTGAGGGGAAATTATAGAGCTTGTGTGACTTTAGGGAGTAGCT 2493
                                                          GCTGTCTATGTGTCCTGGCTGCCCAAGGGACACTCCTGCAGAGCCATTTTTGGGTAAGGA 1961
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RESULT 5
US-09-266-965-51
; Sequence 51, Application US/09266965
; Patent No. 6495348
; GENERAL INFORMATION:
; APPLICANT: Sherman, D
APPLICANT: Varoglu, M
APPLICANT: Sherman, D
APPLICANT: Sheldon, P
TITLE OF INVENTION: Mitomycin biosynthetic gene
; FILE REFERENCE: 600.456US1
; CURRENT APPLICATION NUMBER: US/09/266,965
; CURRENT FILING DATE: 1999-03-12
; EARLIER FILING DATE: 1996-08-19
; EARLIER APPLICATION NUMBER: US 08/624,447
; EARLIER APPLICATION NUMBER: D8/654,447
; EARLIER FILING DATE: 1996-08-19
; EARLIER FILING DATE: 1996-08-19

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                                                                                                                                     ; LENGTH: 53500
; TYPE: DNA
; ORGANISM: Streptomyces lavendulae
US-09-266-965-76
                                                                                                                                                                                                                                                     APPLICANT: VAROGIU, M
APPLICANT: He, M
APPLICANT: Sheldon, P
APPLICANT: Sheldon, P
APPLICANT: Sheldon, P
APPLICANT: Sheldon, P
ITILE OF INVENTION: Mitcomycin blosynthetic get
FILE REFERENCE: 600.456US1
CURRENT APPLICATION NUMBER: US/09/266,965
CURRENT FILING DATE: 1999-03-12
EARLIER APPLICATION NUMBER: US 08/624,447
EARLIER APPLICATION NUMBER: DCT/US94/11279
EARLIER APPLICATION NUMBER: DCT/US94/11279
EARLIER APPLICATION NUMBER: US 08/133,963
EARLIER APPLICATION NUMBER: US 08/133,963
MITMORDEN DO CECT TIN NOC. 116
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US-09-266-965-76
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; ORGANISM: Streptomyces lavendulae
US-09-266-965-51
                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 145
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 76
                                                                 Matches
                                                                                Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Sherman, D
APPLICANT: Mao, Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6495348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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EARLIER FILING DATE: 1993-10-07
NUMBER OF SEQ ID NOS: 145
SOFTWARE: FRETSEQ for Windows Version 3.0
SEQ ID NO 51
LENGTH: 1413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 181;
                                                                              Local Similarity
                                                            3.0%;
cal Similarity 50.8%;
181; Conservative
235 CCCGGAGCGCAGCTACTCGCTGTGCGCGTGACCCCGGACGCCGGCTGCGGGGCCCG 294
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ilarity 50.8%;
Conservative
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                                                      Score 76; DB 4; Length 53500;
Pred. No. 2.8e-07;
0; Mismatches 175; Indels
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Pred. No.
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CURRENT FILING DATE: 2000-07-14
PRIOR APPLICATION NUMBER: US 09/517,849
PRIOR FILING DATE: 2000-03-02
PRIOR APPLICATION NUMBER: US 08/979,608
PRIOR FILING DATE: 1997-11-26
PRIOR APPLICATION NUMBER: US 60/031,930
PRIOR APPLICATION NUMBER: US 60/048,547
PRIOR FILING DATE: 1997-06-03
PRIOR FILING DATE: 1997-06-03
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                                                                                                                                                                                                                                               Query Match 2.8%;
Best Local Similarity 49.1%;
Matches 285; Conservative
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Lees, Ann M.
APPLICANT: Lees, Robert S.
APPLICANT: Lees, Robert S.
APPLICANT: Law, Simon W.
APPLICANT: Arjona, Anibal A.
TITLE OF INVENTION: NOVEL LOW DENSITY LIPOPROTEIN BINDING TITLE OF INVENTION: PROTEINS AND THEIR USE IN DIAGNOSING TITLE OF INVENTION: ATHEROSCLEROSIS
FILE REFERENCE: 10797-004001
                                                                                                                                                                                                                                                                                                                                                FEATURE:
NAME/KEY: CDS
LOCATION: (246)...(1895)
                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Oryctolagus
                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 2561
TYPE: DNA
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1085 GCCGCGGGTCAGGCGGCCAGCGCCGCCGCTACCCCCGAGGTAGCGCACGACTTCCCCG
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                                  CTGGCTCACTTCGCCCTAGGCGCCGACGCCCCCGGCGACGCCGACCCCGGC 214
                                                                               ACGGGTGCGCTCCAGGCGGCCGCCGCCGCCTCCTTCCAGCAGACCCTGCACGCG
                                                                                                                      GCCTTCGCCCGGCTCCTGCGCGGGCCACTGTCGGGGCCGCTGCTCGGGCGGCGCGCGGG
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Pred. No. 6.3e-07;
0; Mismatches 279;
                                                                                                                                                                                                                                                                                      Length 2561;
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; Sequence 1, Application US/09103840A
Patent No. 6294328
; GENERAL INFORMATION:
APPLICANT: FLEISCHMAN, Robert D.
APPLICANT: WHITE, Owen R.
APPLICANT: WHATE, Owen R.
APPLICANT: WHATE, Owen R.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DNA SEQUENCES FOR
TITLE OF INVENTION: TUBERCULOSIS
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US-09-103-840A-1/
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                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: DNA
; ORGANISM: Mycobacterium tuberculosis
; OTHER INFORMATION: H37Rv
US-09-103-840A-1
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CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 47.7
Matches 267; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 4411529
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                                                                2.7%;
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Pred. No. 6.5e-05;
0; Mismatches 291;
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TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN
TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-2007.00
CURRENT APPLICATION NUMBER: US/99/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
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US-09-103-840A-2/c
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Best Local Similarity 47.5%;
Matches 266; Conservative (
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APPLICANT: WHITE, Owen R.
APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, John C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: CDC 1551
OTHER INFORMATION: "n" bases at various positions throughout the sequence
OTHER INFORMATION: represent a, t, c or g
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Pred. No. 0.00016;
0; Mismatches 292;
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                                                                                                           US-08-642-255-60
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GENERAL INFORMATION:
                                                      Matches 268;
                                                                                Query Match
                                                                                                                                                                                       TELEX: 910 277299 FHT UR INFORMATION FOR SEQ ID NO: 60 SEQUENCE CHARACTERISTICS: LENGTH: 810 base pairs
                                                                                                                      TOPOLOGY: line
MOLECULE TYPE: OBSCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: FERRARI, Franco A.
TITLE OF INVENTION: High Molecular Weight Collagen-Like
TITLE OF INVENTION: Protein Polymers
NUMBER OF SEQUENCES: 135
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I.
REGISTRATION NUMBER: 20,015
                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94111-4187

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: A55556-3/BIR TELECOMMUNICATION INFORMATION:
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STATE: California
                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: double
                                                                   Local Similarity
                                                                                                                                                                                                                                               TELEFAX:
                                                                                                                                                                                                                                                            TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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  GGCCCAGCGGGTCCGCCTGGATCTAGAGGTGACCCAGGACCGCCTGGTGCTCCGGGACCT
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(415) 494-8771
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                                                      Conservative
                                                                                                                                                  linear
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                                                                                                                      other nucleic acid
/desc = "synthetic"
                                                                2.7%;
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                                                                   Score 68.2; DB 1;
Pred. No. 2.3e-06;
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                                                      Mismatches 333;
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RESULT 11
US-08-642-255-50
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FILING DATE:
CLASSIFICATION: 435
ATTORNEY_AGENT INFORMATION:
NAME: ROWLAND, Bertram I.
REGISTRATION NUMBER: 20,015
REFERENCE_DOCKET NUMBER: A555
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: CAPPELLO, Joseph
APPLICANT: FERRARI, Franco A.
TITLE OF INVENTION: High Molecular Weight Collagen-Like
TITLE OF INVENTION: Protein Polymers
NUMBER OF SEQUENCES: 135
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/642,25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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RESULT 12
US-09-949-016-15796/C
US-09-949-016-15796, Application US/09949016
Sequence 15796, Application US/09949016
GENERAL INFORMATION:
FAPPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN (
TITLE OF INVENTION: WITH HUMAN DISEASE, MET
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
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INFORMATION FOR SEQ ID NO: 50:
SEQUENCE CHARACTERISTICS:
LENGTH: 756 base pairs
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Pred. No. 4.2e-06;
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US-09-949-016-15796
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FASCSEQ for Windows Version 4.
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LENGTH: 41106
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Pred. No. 5.5e-05;
0; Mismatches 303;
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US-09-902-540-8819/c
(Sequence 8819, Application US/09902540)

Patent No. 6833447
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Sequence 38, Application US/09902540
Patent No. 6833447
GENERAL INFORMATION:
APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION UNMBER: 60/217,883
PRIOR APPLICATION SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERVED SERV
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Best Local Similarity
Matches 285; Conserv
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NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 8819
LENGTH: 6645
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ORGANISM: Myxococcus
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Pred. No. 2.6e-05;
0; Mismatches 312;
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Sequence 193, Application US/10237551

Patent No. 6821519

GENERAL INFORMATION:

APPLICANT: Day, Craig H.

APPLICANT: Hosken, Nancy A.

APPLICANT: Parsons, Joseph M.

TITLE OF INVENTION: TREATMENT OF HERPES SIMPLEX VIRUS INFECTION

TITLE OF INVENTION: TREATMENT OF HERPES SIMPLEX VIRUS INFECTION

FILLE REFERENCE: 210121.538C3

CURRENT APPLICATION NUMBER: US/10/237,551

CURRENT FILING DATE: 2002-09-06

NUMBER OF SEQ ID NOS: 254

SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH: 3957
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US-10-237-551-193
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LENGTH: 10210
TYPE: DNA
ORGANISM: Myxococcus
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Pred. No. 3.3e-05;
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US-10-237-551-193
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